This Raw Listing contains the General Information
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1    <110> APPLICANT: Soto, Ana, et al.
2    <120> TITLE OF INVENTION: A NOVEL ANDROGEN-INDUCED SUPPRESSOR OF CELL PROLIFERATION
3    <130> FILE REFERENCE: MBI-008-1
4    <140> CURRENT APPLICATION NUMBER: US/09/512,581
5    <141> CURRENT FILING DATE: 2000-02-24
6    <160> NUMBER OF SEQ ID NOS: 5
7    <170> SOFTWARE: PatentIn Ver. 2.0
8    <210> SEQ ID NO 1
9    <211> LENGTH: 5271
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24         Val Arg Arg Leu Lys Met Val Val Lys Thr Phe Met Asp Met Asp Gln
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26         gac tct gaa gaa gaa aag gag ctt tat tta aac cta gct tta cat ctt    254
27         Asp Ser Glu Glu Glu Lys Glu Leu Tyr Leu Asn Leu Ala Leu His Leu
28                    50                  55                  60
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30         Ala Ser Asp Phe Phe Leu Lys His Pro Gly Lys Asp Val Arg Leu Leu
31                65                  70                  75
32         gta gcc tgc tgc ctt gct gat att ttc agg att tat gct cct gaa gct    350
33         Val Ala Cys Cys Leu Ala Asp Ile Phe Arg Ile Tyr Ala Pro Glu Ala
34          80                  85                  90                  95
35         cct tac aca tcc cct gat aaa cta aag gat ata ttt atg ttt ata aca    398
36         Pro Tyr Thr Ser Pro Asp Lys Leu Lys Asp Ile Phe Met Phe Ile Thr
37                        100                 105                 110
38         aga cag ttg aag ggg cta gag gat aca aag agc cca caa ttc aat agg    446
39         Arg Gln Leu Lys Gly Leu Glu Asp Thr Lys Ser Pro Gln Phe Asn Arg
40                        115                 120                 125
41         tat ttt tat tta ctt gag aac att gct tgg gtc aag tca tat aac ata    494
42         Tyr Phe Tyr Leu Leu Glu Asn Ile Ala Trp Val Lys Ser Tyr Asn Ile
43                        130                 135                 140
44         tgc ttt gag tta gaa gat agc aat gaa att ttc acc cag cta tac aga    542
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OIPE

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45              Cys Phe Glu Leu Glu Asp Ser Asn Glu Ile Phe Thr Gln Leu Tyr Arg
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47              acc tta ttt tca gtt ata aac aat ggc cac aat cag aaa gtc cat atg    590
48              Thr Leu Phe Ser Val Ile Asn Asn Gly His Asn Gln Lys Val His Met
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50              cac atg gta gac ctt atg agc tct att att tgt gaa ggt gat aca gtg    638
51              His Met Val Asp Leu Met Ser Ser Ile Ile Cys Glu Gly Asp Thr Val
52                              180             185             190
53              tct cag gag ctt ttg gat acg gtt tta gta aat ctg gta cct gct cat    686
54              Ser Gln Glu Leu Leu Asp Thr Val Leu Val Asn Leu Val Pro Ala His
55                              195             200             205
56              aag aat tta aac aag caa gca tat gat ttg gca aag gct tta ctg aag    734
57              Lys Asn Leu Asn Lys Gln Ala Tyr Asp Leu Ala Lys Ala Leu Leu Lys
58                          210             215             220
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60              Arg Thr Ala Gln Ala Ile Glu Pro Tyr Ile Thr Thr Phe Phe Asn Gln
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62              gtt ctg atg ctt ggg aaa aca tct atc agc gat ttg tca gag cat gtc    830
63              Val Leu Met Leu Gly Lys Thr Ser Ile Ser Asp Leu Ser Glu His Val
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69              Ser Val Leu Pro Gln Leu Glu Phe Lys Leu Lys Ser Asn Asp Asn Glu
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71              gag cgc cta caa gtt gtt aaa cta ctg gca aaa atg ttt ggg gca aag    974
72              Glu Arg Leu Gln Val Val Lys Leu Leu Ala Lys Met Phe Gly Ala Lys
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74              gat tca gaa ttg gct tct caa aac aag cca ctt tgg cag tgc tac ttg   1022
75              Asp Ser Glu Leu Ala Ser Gln Asn Lys Pro Leu Trp Gln Cys Tyr Leu
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77              ggc agg ttt aat gat atc cat gta cca atc cgc ctg gaa tgt gtg aaa   1070
78              Gly Arg Phe Asn Asp Ile His Val Pro Ile Arg Leu Glu Cys Val Lys
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81              Phe Ala Ser His Cys Leu Met Asn His Pro Asp Leu Ala Lys Asp Leu
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84              Thr Glu Tyr Leu Lys Val Arg Ser His Asp Pro Glu Glu Ala Ile Arg
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86              cat gat gtt att gtg tca ata gtt aca gct gct aaa aag gat att ctt   1214
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90              Leu Val Asn Asp His Leu Leu Asn Phe Val Arg Glu Arg Thr Leu Asp
91                          385             390             395
92              aaa cga tgg aga gta cgc aaa gaa gcc atg atg gga ctt gcc caa att   1310
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101        agt att gat gat cga cta ctt gtt gaa cgg atc ttt gct caa tac atg    1454
102        Ser Ile Asp Asp Arg Leu Leu Val Glu Arg Ile Phe Ala Gln Tyr Met
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104        gtt cct cac aat tta gaa act aca gaa cgg atg aaa tgc tta tat tac    1502
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122        aga aag cag tta gaa gta ctt gtt agt cca aca tgc tcc tgc aag cag    1790
123        Arg Lys Gln Leu Glu Val Leu Val Ser Pro Thr Cys Ser Cys Lys Gln
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125        gct gaa ggt tgt gtg cgt gaa ata act aag aag ttg ggc aac ccc aaa    1838
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141        Leu Ser Phe Thr His Pro Ile Ser Phe His Ser Ala Glu Thr Phe Glu
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143        tca tta ctg gct tgt ctg aaa atg gat gat gaa aaa gta gca gaa gct    2126
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153	Lys	Gly	Pro	Pro	Arg	Gln	Ala	Lys	Tyr	Ala	Ile	His	Cys	Ile	His	Ala	
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156	Ile	Phe	Ser	Ser	Lys	Glu	Thr	Gln	Phe	Ala	Gln	Ile	Phe	Glu	Pro	Leu	
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176	act	tct	acc	tta	aga	ttg	cta	aca	aca	ata	ttg	cat	agt	gat	gga	gac	2654
177	Thr	Ser	Thr	Leu	Arg	Leu	Leu	Thr	Thr	Ile	Leu	His	Ser	Asp	Gly	Asp	
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180	Leu	Thr	Glu	Gln	Gly	Lys	Ile	Ser	Lys	Pro	Asp	Met	Ser	Arg	Leu	Arg	
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185	cat	gaa	atc	atc	aca	tta	gaa	caa	tat	cag	cta	tgt	gca	tta	gct	atc	2798
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212         atc aga aag atg gta gaa aat att aaa caa aca aaa gat gcc caa gga    3230
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